



ENTERED

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,014

DATE: 03/05/2002

TIME: 14:12:21

Input Set : A:\sequence list.txt

Output Set: N:\CRF3\03052002\I701014.raw

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3 <110> APPLICANT: Mitraphanous, Kyrlacos
4     Patil, Deva
5     Kingsman, Alan
6     Kingsman, Susan
7     Ellard, Fiona
9 <120> TITLE OF INVENTION: Retroviral Delivery System
11 <130> FILE REFERENCE: 550-184
13 <140> CURRENT APPLICATION NUMBER: 09/701,014
14 <141> CURRENT FILING DATE: 2001-05-24
16 <150> PRIOR APPLICATION NUMBER: PCT/GB99/01607
17 <151> PRIOR FILING DATE: 1999-05-21
19 <150> PRIOR APPLICATION NUMBER: GB 9811153.7
20 <151> PRIOR FILING DATE: 1998-07-17
22 <160> NUMBER OF SEQ ID NOS: 4
24 <170> SOFTWARE: PatentIn Ver. 2.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 1650
28 <212> TYPE: DNA
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Description of Artificial Sequence: rabies virus strain ERA
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36 tgggaaattc cctattttaca cgatactaga caagcttggt ccctggagcc cgattgacat 120
37 acatcacctc agctgcccaa acaatttggt agtggaggac gaaggatgca ccaacctgtc 180
38 agggttctcc tacatggaac ttaaagttgg atacatctta gccataaaaa tgaacgggtt 240
39 cacttgcaca ggcgttgtga cggaggctga aacctacact aacttcgttg gttatgtcac 300
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45 gcccgagaat ccgagactag ggatgtcttg tgacattttt accaatagta gagggaagag 660
46 agcatccaaa gggagtgaga cttgcggctt tgtagatgaa agaggcctat ataagtcttt 720
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58 gtatgtatta ctgagtgcag gggccctgac tgccttgatg ttgataattt tcctgatgac 1440
59 atgttgtaga agagtcaatc gatcagaacc tacgcaacac aatctcagag ggacagggag 1500
60 ggaggtgtca gtcactcccc aaagcgggaa gatcatatct tcatgggaat cacacaagag 1560
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77 20 25 30
79 Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val
80 35 40 45
82 Val Glu Asp Glu Gly Cys Thr Asn Leu Ser Gly Phe Ser Tyr Met Glu
83 50 55 60
85 Leu Lys Val Gly Tyr Ile Leu Ala Ile Lys Met Asn Gly Phe Thr Cys
86 65 70 75 80
88 Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
89 85 90 95
91 Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Pro Asp Ala
92 100 105 110
94 Cys Arg Ala Ala Tyr Asn Trp Lys Met Ala Gly Asp Pro Arg Tyr Glu
95 115 120 125
97 Glu Ser Leu His Asn Pro Tyr Pro Asp Tyr Arg Trp Leu Arg Thr Val
98 130 135 140
100 Lys Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Val Ala Asp
101 145 150 155 160
103 Leu Asp Pro Tyr Asp Arg Ser Leu His Ser Arg Val Phe Pro Ser Gly
104 165 170 175
106 Lys Cys Ser Gly Val Ala Val Ser Ser Thr Tyr Cys Ser Thr Asn His
107 180 185 190
109 Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Arg Leu Gly Met Ser Cys
110 195 200 205
112 Asp Ile Phe Thr Asn Ser Arg Gly Lys Arg Ala Ser Lys Gly Ser Glu
113 210 215 220
115 Thr Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly
116 225 230 235 240
118 Ala Cys Lys Leu Lys Leu Cys Gly Val Leu Gly Leu Arg Leu Met Asp
119 245 250 255
121 Gly Thr Trp Val Ala Met Gln Thr Ser Asn Glu Thr Lys Trp Cys Pro
122 260 265 270
124 Pro Asp Gln Leu Val Asn Leu His Asp Phe Arg Ser Asp Glu Ile Glu

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125 275 280 285
 127 His Leu Val Val Glu Glu Leu Val Arg Lys Arg Glu Glu Cys Leu Asp
 128 290 295 300
 130 Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu
 131 305 310 315 320
 133 Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile
 134 325 330 335
 136 Phe Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg
 137 340 345 350
 139 Thr Trp Asn Glu Ile Leu Pro Ser Lys Gly Cys Leu Arg Val Gly Gly
 140 355 360 365
 142 Arg Cys His Pro His Val Asn Gly Val Phe Phe Asn Gly Ile Ile Leu
 143 370 375 380
 145 Gly Pro Asp Gly Asn Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu
 146 385 390 395 400
 148 Gln Gln His Met Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Val His
 149 405 410 415
 151 Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Asp Gly Asp Glu Ala Glu
 152 420 425 430
 154 Asp Phe Val Glu Val His Leu Pro Asp Val His Asn Gln Val Ser Gly
 155 435 440 445
 157 Val Asp Leu Gly Leu Pro Asn Trp Gly Lys Tyr Val Leu Leu Ser Ala
 158 450 455 460
 160 Gly Ala Leu Thr Ala Leu Met Leu Ile Ile Phe Leu Met Thr Cys Cys
 161 465 470 475 480
 163 Arg Arg Val Asn Arg Ser Glu Pro Thr Gln His Asn Leu Arg Gly Thr
 164 485 490 495
 166 Gly Arg Glu Val Ser Val Thr Pro Gln Ser Gly Lys Ile Ile Ser Ser
 167 500 505 510
 169 Trp Glu Ser His Lys Ser Gly Gly Glu Thr Arg Leu
 170 515 520
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 174 <211> LENGTH: 44
 175 <212> TYPE: DNA
 176 <213> ORGANISM: Artificial Sequence
 178 <220> FEATURE:
 179 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
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 185 <211> LENGTH: 44
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 187 <213> ORGANISM: Artificial Sequence
 189 <220> FEATURE:
 190 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
 192 <400> SEQUENCE: 4
 193 gggaggatct cattccaagt ctggactgac ttgtagtgag catc 44

VERIFICATION SUMMARY

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date

09/01/04 14:12:22